- SEQUENCE LISTING

<110> Hoechst Marion Roussel

<120> MATURE PROTEIN HAVING ANTAGONIST ACTIVITY AGAINST BONE MORPHOGENETIC PROTEIN.

<130> JH98K011 PCT SEQUENCES IN ENGLISH

<140>

<141>

<150> 10-288103

<151> 1998-10-09

<160> 7

<170> PatentIn Ver. 2.1

<210> 1

<211> 119

<212> PRT

<213> Human

<220>

<221> CHAIN

<222> (1)..(119)

<223> Mature MP52

<300>

<301> MAKISHIMA, Fusoa

TAKAMATSU, Hiroyuki

MIKI, Hideo

KAWAI, Shinji

KIMURA, Michio

MATSUMOTO, Tomoaki

KATSUURA, Mieko

ENOMOTO, Koichi

SATOH, Yusuke

<302> Novel protein and process for producing the same.

<310> WO 96/33215

<312> 1996-10-24

<313> 1 TO 119

<400> 1

Pro Ser Ala Thr Arg Gln Gly Lys Arg Pro Ser Lys Asn Leu Lys Ala

Arg Cys Ser Arg Lys Ala Leu His Val Asn Phe Lys Asp Met Gly Trp \$20\$

Asp Asp Trp Ile Ile Ala Pro Leu Glu Tyr Glu Ala Phe His Cys Glu 35 40 45

Gly Leu Cys Glu Phe Pro Leu Arg Ser His Leu Glu Pro Thr Asn His 50 55 60

Ala Val Ile Gln Thr Leu Met Asn Ser Met Asp Pro Glu Ser Thr Pro 65 70 75 80

Pro Thr Cys Cys Val Pro Thr Arg Leu Ser Pro Ile Ser Ile Leu Phe 85 90 95

Ile Asp Ser Ala Asn Asn Val Val Tyr Lys Gln Tyr Glu Asp Met Val $100 \hspace{1.5cm} 105 \hspace{1.5cm} 110$

Val Glu Ser Cys Gly Cys Arg

<210> 2

<211> 114

<212> PRT

<213> Human

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<221> CHAIN

<222> (1) .. (114)

<223> Mature BMP-2

<300>

<301> WANG, Elizabeth A.

WOZNEY, John M.

ROSEN, Vicki A.

<302> Novel osteoinductive compositions.

<310> WO 88/00205

<312> 1988-01-14

<313> 1 TO 114

<400> 2

Gln Ala Lys His Lys Gln Arg Lys Arg Leu Lys Ser Ser Cys Lys Arg

His Pro Leu Tyr Val Asp Phe Ser Asp Val Gly Trp Asn Asp Trp Ile 20 25 30

Val Ala Pro Pro Gly Tyr His Ala Phe Tyr Cys His Gly Glu Cys Pro 35 40 45

Phe Pro Leu Ala Asp His Leu Asn Ser Thr Asn His Ala Ile Val Gln

Thr Leu Val Asn Ser Val Asn Ser Lys Ile Pro Lys Ala Cys Cys Val 65 70 75 80

Pro Thr Glu Leu Ser Ala Ile Ser Met Leu Tyr Leu Asp Glu Asn Glu 85 90 95

Lys Val Val Leu Lys Asn Tyr Gln Asp Met Val Val Glu Gly Cys Gly 100 105 110

' WO 00/21998 ' PCT/IB99/01621

<210> 3

<211> 116

<212> PRT

<213> Human

<220>

<221> CHAIN

<222> (1)..(116)

<223> Mature BMP-4

<300>

<301> WOZNEY, John M.

ROSEN, Vicki

CELESTE, Anthony J.

MITSOCK, Lisa M.

WHITTERS, Matthew J.

KRIZ, Ronald W.

HEWICK, Rodney M.

WANG, Elizabeth A.

<302> Novel regulators of bone formation : molecular clones and activities.

<303> Science

<304> 242

<305> 4885

<306> 1528-1534

<307> 1988-12-16

<308> Genbank/M22490

<313> 1 TO 116

<400> 3

Ser Pro Lys His His Ser Gln Arg Ala Arg Lys Lys Asn Lys Asn Cys

Arg Arg His Ser Leu Tyr Val Asp Phe Ser Asp Val Gly Trp Asn Asp $20 \hspace{1.5cm} 25 \hspace{1.5cm} 30$

Trp Ile Val Ala Pro Pro Gly Tyr Gln Ala Phe Tyr Cys His Gly Asp

40

Cys Pro Phe Pro Leu Ala Asp His Leu Asn Ser Thr Asn His Ala Ile 55 50

Val Gln Thr Leu Val Asn Ser Val Asn Ser Ser Ile Pro Lys Ala Cys 75 70 65

Cys Val Pro Thr Glu Leu Ser Ala Ile Ser Met Leu Tyr Leu Asp Glu 90 85

Tyr Asp Lys Val Val Leu Lys Asn Tyr Gln Glu Met Val Val Glu Gly 105 100

Cys Gly Cys Arg 115

<210> 4

<211> 139

<212> PRT

<213> Human

<220>

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<221> CHAIN

<222> (1)..(139)

<223> Mature BMP-7

<300>

<301> OZKAYNAK, Engin

RUEGER, David C.

DRIER, Eric A.

CORBETT, Clare

RIDGE, Richard J.

SAMPATH, Kuber T.

OPPERMANN, Hermann

<302> OP-1 cDNA encodes an osteogenic protein in the TGF-beta family.

<303> EMBO J. <304> 9 <305> 7 <306> 2085-2093 <307> 1990 <308> EMBL data library/X51801 <313> 1 TO 139

<400> 4

Ser Thr Gly Ser Lys Gln Arg Ser Gln Asn Arg Ser Lys Thr Pro Lys 10

Asn Gln Glu Ala Leu Arg Met Ala Asn Val Ala Glu Asn Ser Ser Ser 25 20

Asp Gln Arg Gln Ala Cys Lys Lys His Glu Leu Tyr Val Ser Phe Arg 35

Asp Leu Gly Trp Gln Asp Trp Ile Ile Ala Pro Glu Gly Tyr Ala Ala 55 50

Tyr Tyr Cys Glu Gly Glu Cys Ala Phe Pro Leu Asn Ser Tyr Met Asn 75 70 65

Ala Thr Asn His Ala Ile Val Gln Thr Leu Val His Phe Ile Asn Pro 90 85

Glu Thr Val Pro Lys Pro Cys Cys Ala Pro Thr Gln Leu Asn Ala Ile 105 100

Ser Val Leu Tyr Phe Asp Asp Ser Ser Asn Val Ile Leu Lys Lys Tyr 120 115

Arg Asn Met Val Val Arg Ala Cys Gly Cys His 135 130

PCT/IB99/01621

· WO 00/21998 7

<210> 5

<211> 119 <212> PRT

<213> Human

<220>

<221> CHAIN

<222> (1)..(119)

<223> Mature MP52 protein. Note : 30th, 71st, 74th and 111th Met are modified to Met sulfoxide.

Tergi

Pro Ser Ala Thr Arg Gln Gly Lys Arg Pro Ser Lys Asn Leu Lys Ala <400> 5 10 5 1

Arg Cys Ser Arg Lys Ala Leu His Val Asn Phe Lys Asp Met Gly Trp 20

Asp Asp Trp Ile Ile Ala Pro Leu Glu Tyr Glu Ala Phe His Cys Glu 40 35

Gly Leu Cys Glu Phe Pro Leu Arg Ser His Leu Glu Pro Thr Asn His 60 55 50

Ala Val Ile Gln Thr Leu Met Asn Ser Met Asp Pro Glu Ser Thr Pro 75 70 65

Pro Thr Cys Cys Val Pro Thr Arg Leu Ser Pro Ile Ser Ile Leu Phe 90

Ile Asp Ser Ala Asn Asn Val Val Tyr Lys Gln Tyr Glu Asp Met Val 105 100

Val Glu Ser Cys Gly Cys Arg

<210> 6 <211> 119

<212> PRT

<213> Human

<220>

<221> CHAIN

<222> (1)..(119)

<223> Mature MP52 protein. Note : 30th and/or 71st and/or 74th and/or 111th Met are modified to s-carboxymethyl Met.

<400> 6

W.

Pro Ser Ala Thr Arg Gln Gly Lys Arg Pro Ser Lys Asn Leu Lys Ala 15 10 5 1

Arg Cys Ser Arg Lys Ala Leu His Val Asn Phe Lys Asp Met Gly Trp 30 25 20

Asp Asp Trp Ile Ile Ala Pro Leu Glu Tyr Glu Ala Phe His Cys Glu 45 40 35

Gly Leu Cys Glu Phe Pro Leu Arg Ser His Leu Glu Pro Thr Asn His 60 55 50

Ala Val Ile Gln Thr Leu Met Asn Ser Met Asp Pro Glu Ser Thr Pro 75 70 65

Pro Thr Cys Cys Val Pro Thr Arg Leu Ser Pro Ile Ser Ile Leu Phe 90 85

Ile Asp Ser Ala Asn Asn Val Val Tyr Lys Gln Tyr Glu Asp Met Val 110 105 100

Val Glu Ser Cys Gly Cys Arg

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<210> 7

<211> 119

<213> Human

<220>

<221> CHAIN

<222> (1)..(119)

<223> Mature MP52 protein. Note : 32nd and 35th Trp are modified to allylsulphenyl Trp.

<400> 7

Pro Ser Ala Thr Arg Gln Gly Lys Arg Pro Ser Lys Asn Leu Lys Ala

Arg Cys Ser Arg Lys Ala Leu His Val Asn Phe Lys Asp Met Gly Trp 20 25 30

Asp Asp Trp Ile Ile Ala Pro Leu Glu Tyr Glu Ala Phe His Cys Glu 35 40 45

Gly Leu Cys Glu Phe Pro Leu Arg Ser His Leu Glu Pro Thr Asn His

Ala Val Ile Gln Thr Leu Met Asn Ser Met Asp Pro Glu Ser Thr Pro 65 70 75 80

Pro Thr Cys Cys Val Pro Thr Arg Leu Ser Pro Ile Ser Ile Leu Phe 85 90 95

Ile Asp Ser Ala Asn Asn Val Val Tyr Lys Gln Tyr Glu Asp Met Val

Val Glu Ser Cys Gly Cys Arg